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Variation at Isozyme Loci in Phaseolus vulgaris

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Electrophoresis of raw extracts with subsequent staining for specific enzyme activities has been used to measure variability in numerous plant species. In <u>Phaseolus vulgaris</u> distinct allozymes have been described in several isozyme systems (1-3). These studies demonstrated that allozymic variation is present among the major classes of cultivated beans and in closely related species. Unfortunately, only a small number of assays were used in the surveys, severely restricting the amount of variability that could be identified. A compilation of the polymorphic isozyme loci in this species would be very useful for cultivar identification, marking of commercially important traits and further genetic characterization of the bean genome. With such a goal in mind I have initiated a survey of isozyme variability in this species and report here the identification of polymorphism at 15 loci during a survey of 134 lines using 27 different assays.

Electrophoresis was performed on horizontal starch gels using methods and assays described in Weeden and Gottlieb (4), Shaw and Prasad (5) and Weeden (in prep.). Seeds were obtained from Dr. M. P. Dickson at the N.Y. State Agricultural Experiment Station or directly from seed companies. An intense effort was made in snap beans to identify loci which would be useful for distinguishing cultivars. Each cultivar was then screened at most if not all these loci to determine their genotype. In most cases over 20 plants were tested from each line. Additional bean samples, representing other commercial classes or experimental lines, were also subjected to electrophoretic analysis using a smaller number of assays.

The results of this survey are presented in Tables 1 and 2. Table 1 consists of the snap bean varieties and breeding lines surveyed. These are listed in a sequence based on the basis of isozyme fingerprints, placing cultivars with identical fingerprints adjacent to each other. Gccasionally, more than one isozyme fingerprint was found to characterize a particular cultivar and such variation is indicated by a second entry for that cultivar in the table. Table 2 lists the additional varieties and lines primarily on the basis of their commercial class.

The snap bean cultivars exhibited a minimum of 79 distinct fingerprints out of a possible 6912 different combinations. Fifty-five of the 103 cultivars possessed unique fingerprints and, thus, could be easily and reliably identified as seedlings. Several lines had to be described by two fingerprints, this ambiguity being caused by allelic variants at one locus. However, this variation did not seriously affect the identification of cultivars. Certain other cultivars, such as Bountiful and Roma or Black Valentine and Pure Gold, which could not be distinguished by isozyme fingerprint are easily differentiated using morphological characters.

In addition to the loci listed in the tables, 6 other loci were found to be polymorphic: glucose phosphate isomerase, xanthine dehydrogenase, succinate dehydrogenase, malate dehydrogenase, isocitrate dehydrogenase and

amylase. Many of the less common alleles were found only in the lines listed in Table 2. This latter group, as might be anticipated, exhibited greater variability in isozyme phenotype than did the snap beans. However, both the xanthine dehydrogenase and succinate dehydrogenase phenotypes differed among snap bean lines and may aid in the differentiation of cultivars.

## References

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- (3) Bassiri, A. and M.W. Adams 1978. An electrophoretic survey of seedling isozymes in several <u>Phaseolus</u> species. Euphytica 27:447-459.
- (4) Weeden, N.F. and L.D. Gottlieb 1980. Isolation of cytoplasmic enzymes from pollen. Plant Physiol. 66:400-403.
- (5) Shaw, C.R. and R. Prasad 1970. Starch gel electrophoresis of enzymes-a compilation of recipes. Biochem. Genet. 4:297-320.

TABLE 1

VARIETY	OBSERVED ALLOZYMES 1,2	VARIETY	OBSERVED ALLOZYMES 1,2
	1 2 3 4 5 6 7 8 9 10		1 2 3 4 5 6 7 8 9 10
FLO	FFFFFFSFS-	BOUNTIFUL	S S F F F S F F F
	FFFFFSFSS	ROMA	SSFFFSFFF-
LAKELAND	FFFFSFFSS	CASCADE	SSFFFSFFSS
TENDERPOD	FFFFSSFSS-	EAGLE	SSFFFSFFSS
LANCER		GOLDRUSH	SSFFFSFFSS
KENTUCKY WONDER	. , 5 , 5 , 5 ,	NCX 8014	SSFFFSFFSS
EARLY BIRD	FFSFFSSFS-	FEAK	SSFFFSFFSS
BBF - 1 4	FFSFSFF#SSF		SSFFFSFFSS
UREGAN 55	FFSFSFF*SSS	XBF 108	
OREGAN 83	FFSFSFF*SSS	BONANZA	
80BF-65	F F S F S S F S S F	SPRITE	SSFFFSFS-
80BP-70	FFSFSSFSS.	STRIKE	S S F F F S S S S -
30BF-84	FFSFSSFSS	TENDERLAKE	SSFFSFS+FS-
8 BF: - 1 4	FFSFSSF*SSF	BBL-GV2	SSFFSSSFS-
ORBIT	FFSSSSFFS-	EXPRESSWAY	SSFFSSSFS-
EPOCH	FFSSSF*SS=	BLACK VALENTINE	SSFSFS-FS-
AURORA	FFUFSUSSSF	FURE GOLD	S
MCCASLAN	FF+SFSSSS ~	68953	S
DANDY	FSFFFFFFS-	PROVIDER	SSFSFSF5-
FIROL	FSFFSFFSS	GOLDEN POD	SSFSFSFSS
STRETCH	FSFSSSFFS-	GREEN CROP	SSFSFSFSS
	FSFSSSFF-	REBEL	SSFSFSFSS
GREENSLEEVES	F S F S S S F F -	W26	SSFSFSFSS
MIAMI		IMPROVED TOPNOTCH	SSFSFSFSS-
PIROL	· · · · · · · · · · · · · · · · · · ·	PLATEAU PROVIDER	SSFSFSFSS-
GREENWAY	rasaraiaa	SPLENDOR GOLD	S S F S F S F S S -
SLIMGREEN	F S S S F S S S S =	EARLIWAX	SSFSFSSFSS
SPARTAN PRIDE	SFFFFFSFS -	RODEO	SSFSFSSFSS
COMETA	SFFFFFSS =		SSFSFSSFSS
TENDERCROP	SFFFFS-SS-	SINCLAIR BUTTERWAX	
TENDERETTE	S F F F S - S S -	MIDAS	
WIN .	SF.FFSFFS-	RAIDER	3 0 , 0 , 2 - ,
BBL-109	SFFFFSFSS	GOLDEN WAX	
EARLY GALLATIN	SFFFFSFSSS	GOLD CROP	SSFSSFFSS
G4219	SFFFFSFSSS	DEL RAY	S S F S S S S - S S
GALLATIN 50	.SFFFFSFSSS	CUMBERLAND	SSFSSSFSS
PAYMASTER	SFFFFSFSS	HARVESTER	SSFSSSSF <b>S</b> S
SLENDERETTE	SFFFFSFSS	SLENDERWHITE	S S F S S S S S -
WONDERGREEN	SFFFFSSSS	MOONGOLD	SSSFFSFFSS
GRAND CANYON	S F F F S S* S S	8BF-8	SSSFFSFSS-
SPARTAN PRIDE	SFFFSFSFS-	REGAL	S
GALASLIN	SFFFSSFSS-	GATORGREEN	S S S F F S S S S -
VALIANT	SFFFSSSFS-	LAKE GENEVA	SSSFSFFFSS
VITAGREEN	S F F F S S S* S S -	8BL-92	SSSFSSFFSS
GINA	SFFSFSFFF-	LAKE SUPERIOR	SSSFSSFSS
MAJESTIC	SFFSFSF5-	IMPROVED TENDERGREEN	SSSSFSFFS-
SUNRISE	S F F S S S S * F S -	GOLD COAST	SSSSFSFSF-
CHECKNATE	SFSFFSFS-	BURLY	SSSSFSFSS-
GREEN ISLE	SFSFFSFSS	IMPROVED HIGRADE	SSSSFSFSS-
SLENDERETTE	SFSFFSFSSS	REGAL	SSSFSSFSS
	S F S F F S F* F S -	BBL-274	SSSSSFFFS
BBL-47		BLUE CROP	SSSSSFFFS
LAKE SHASTA			S S S S S F S S -
SHASTA		BURLY	S S S S S S F F S
OREGAN 43	SFSFSFF*SS-	BBL - 274	
OREGAN 91G	SFSFSFSS =	LAKE SENECA	
GREENPAK	SFSSFSS ~	G742	
BBL-283	S F S S S F* S S -	TORRENT .	S S S S S S S S -

<sup>1</sup> Columns represent 10 polymorphic isozyme loci. The sequence of isozymes is: (1) shikimic dehydrogenase, (2) malic enzyme, (3) diaphorase, (4) ribulose bisphosphate carboxylase, (5) adenylate kinase, (6) esterase-1, (7) esterase-2, (8) peptidase, (9) acid phosphatase, (10) cathodal peroxidase.

Genotype for each cultivar is given for each locus by a single letter abbreviation of the isozyme phenotype observed. Abbreviations are: S = slow;  $S^* = low$  activity of slow allozyme; F = fast;  $F^* = low$  activity of fast allozyme;  $F^* = allozyme$  with mobility faster than F; U = different than S, F, or  $F^*$ ; dash F no data.

Table 2

Variety		_				Allozymes			_	1.0
	1	2	3	4	5	6	7	8	9	10
<u>Dry beans</u>										
Brazil 2	-	-	-	F	F	U	S	-	_	S
Ciat P560	F	F+	U	F	-	U	S	_	S	-
Horsehead	S	S	F	F	F	F	-	F	F	_
State Half Runner	F	F	S	F	S	S	S	S	S	F
Swedish Brown	S	-	-	-	F	-	F	-	-	-
Green shell										
Genuine Cornfield	F	F	F	F	S	_	U	F	S	-
Limelight	F	F+	S	-	S	F	-	F	s	-
Navy (pea)										
Ex Rico	_	_	_	F	S	_	S	_	_	_
Sanalac	F	F+	S	_	S	_	_	S	S	_
Seafarer	F	F+	Š	F	Š	_	_	S	S	_
3322 21 31	-		_	_						
Red Kidney										
California RK	S	S	F	S	F	S	F	F	-	
Cal LRK	-	-	F	S	-	-	F	-	S	_
Red Kloud	S	S	F	S	-	-	F	-	S	_
Red Kloud GC	-	F	F	-	F	-	F	_	S	-
Sacramento Red	S	S	F	S	-	-	F	-	S	_
Other										
336	S	_	_	S	F	Ü	S	_	_	S
924R	_	_	_	F	F	_	F	_	-	S
2114-12	F	U	S	F	S	Ü	S	_	S	S
9505	_	_	_	F	F	Ū	F	_	_	S
165426 c	_	F	S	F	S	Ü	S	_	S	S
165 426 w	-	F	S	F	_	Ü	S	_	S	-
165435	_	F	S	F	S	U	S	_	S	_
Andino	S	Ü	F	S	S	_	-	F	F	_
G06 391	-	Ü	F	F	F	_	F	F	S	-
G08050	_	U	F	S	_	-	F	_	-	_
ICA Tui	F	_	_	-	S	_	S	_	-	S
L1	S	S	S	_	-	-	S	-	S	-
P459	F	-	_	_	S	_	S	-	-	S
P717	F	F+	U	F	S	-	S#	S	S	F
Porillo Sentetica	F	-	-	-	S	U	F	-	S	S
Remus	F	S	F	-	F	-	F	-	-	-

Column and letter designations are the same as for Table 1.